

FIG. 1

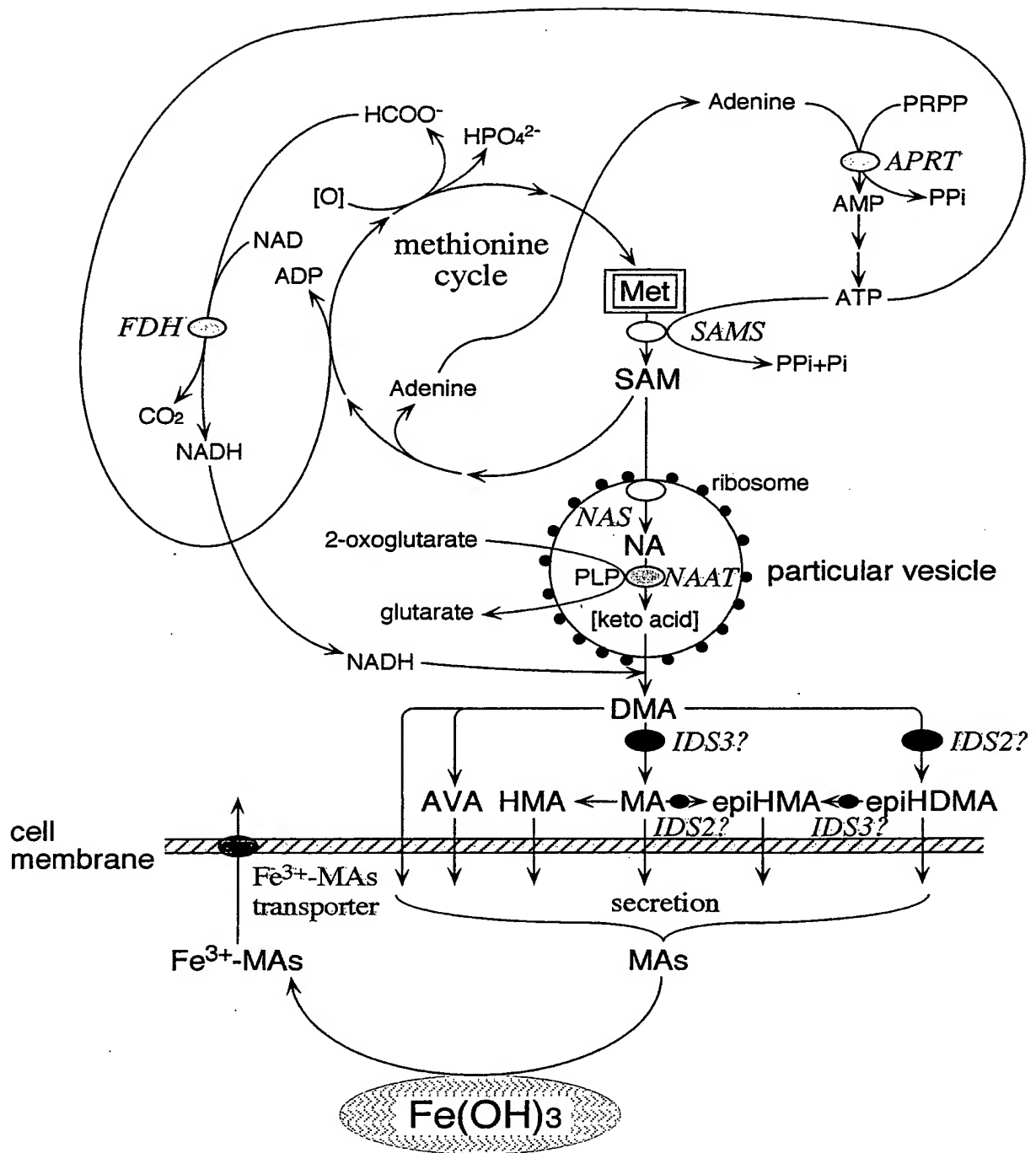
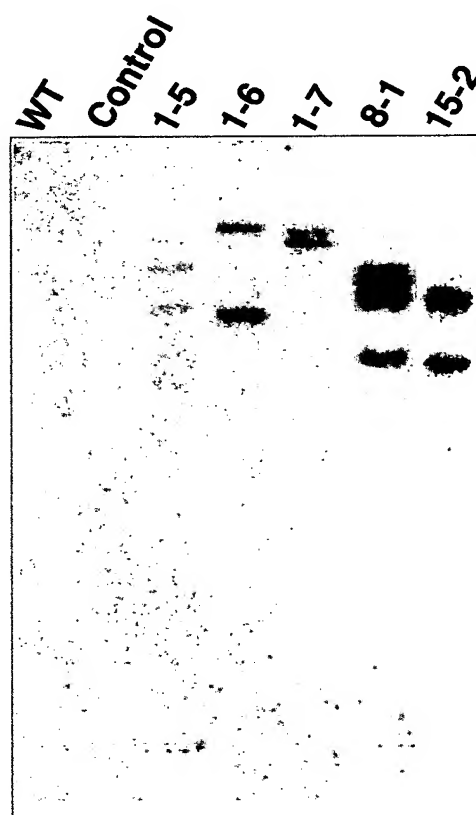


FIG. 2



FIG. 3



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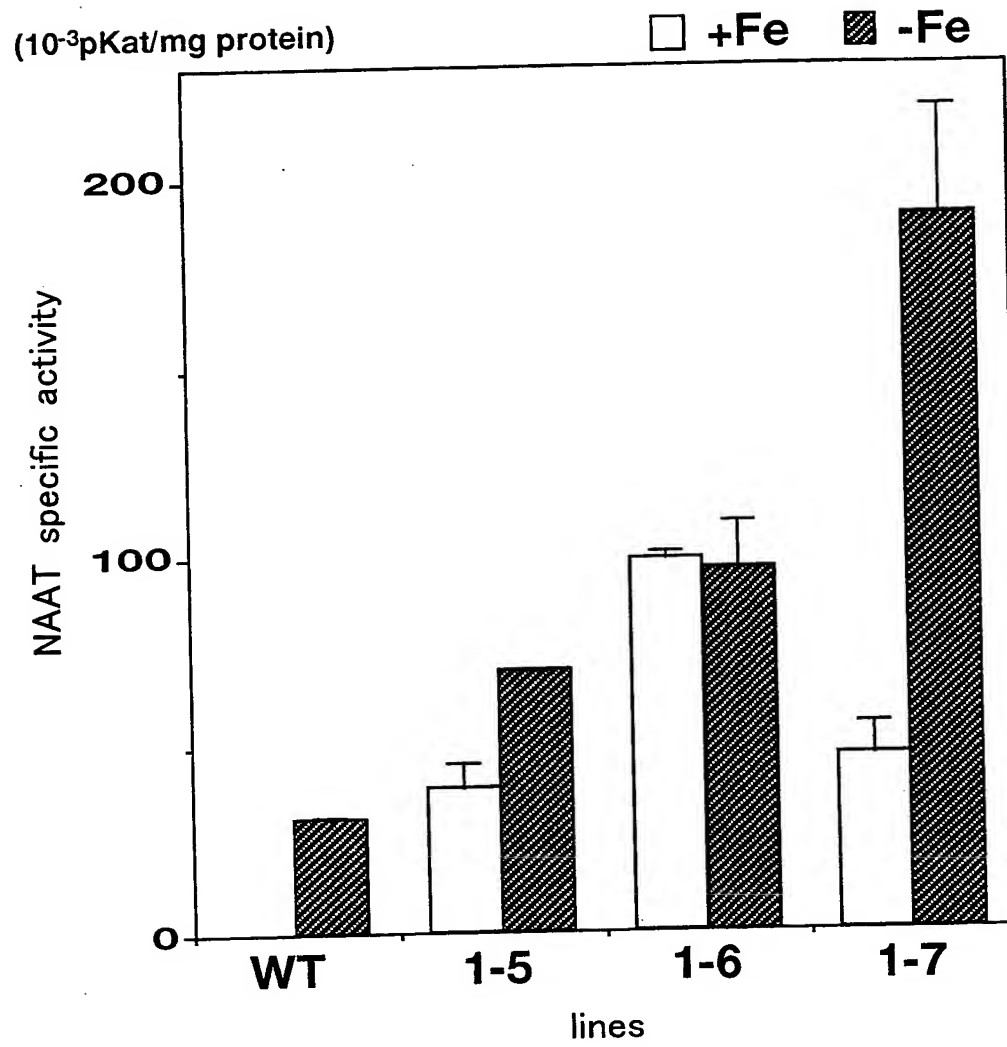


FIG. 5



FIG. 6

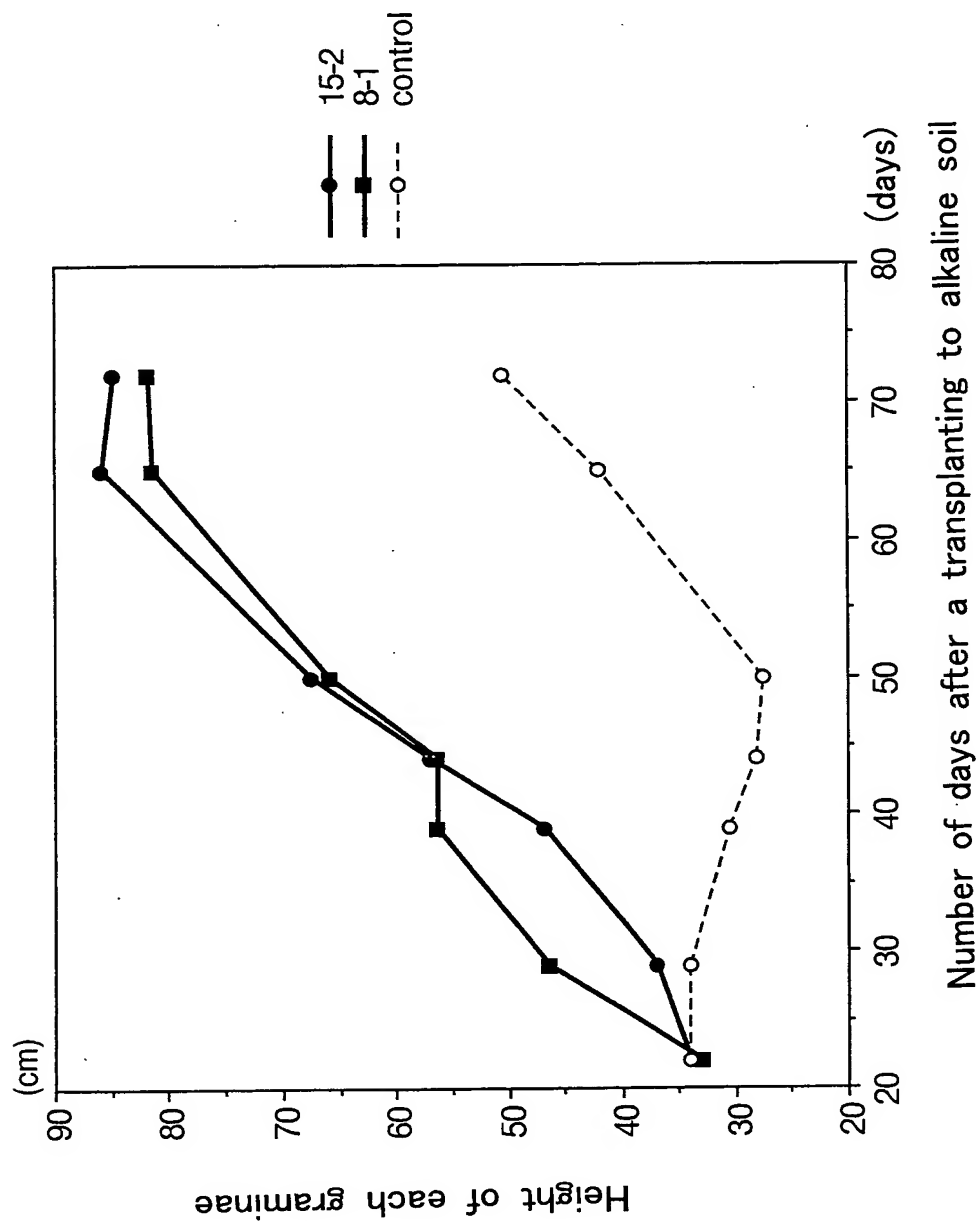
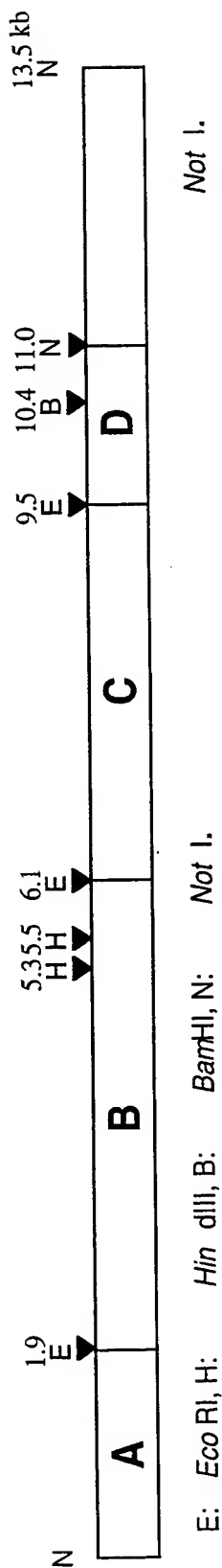
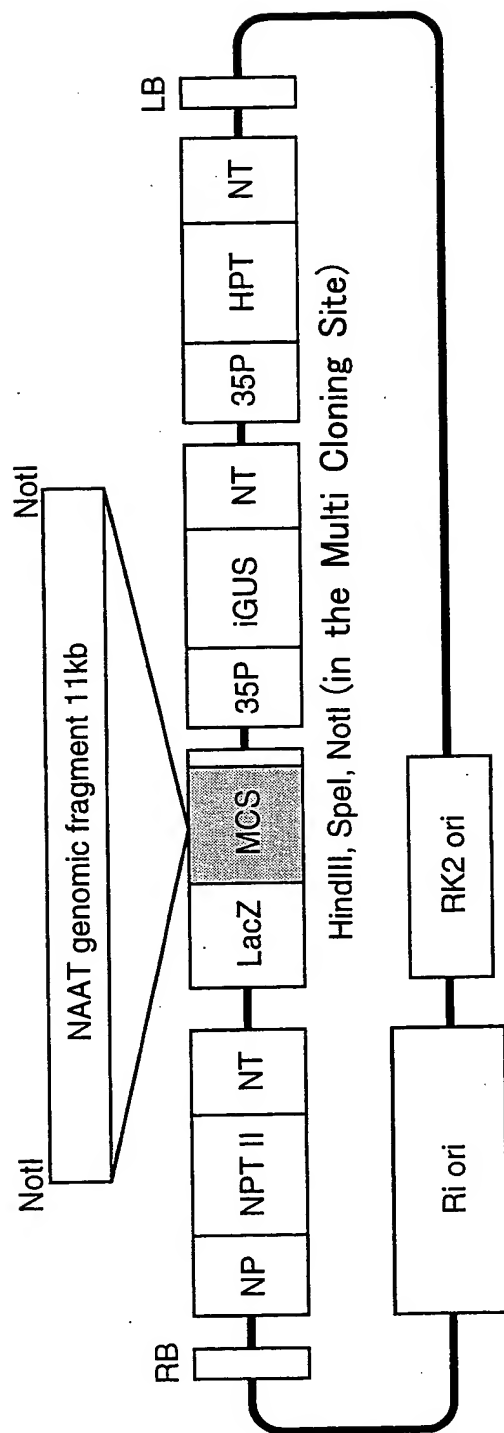


FIG. 7



The Not I site on both sides is the Not I located at the arm of λFIX II

FIG. 8



F I G . 9

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 M V H Q S N G H NAAT-A

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 G E A A A A A A N G K S N G H A A A A N

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 D G I L A T T G A K N S I R A I R Y K I

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 S A S V E E S G P R P V L P L A H G D P

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 S V F P A F R T A V E A E D A V A A A L

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 R T G Q F N C Y A A G V G L P A A R S

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Q T A G A N I L L P R P G Y P N Y E A R

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A A F N K L E V R H F D L I P D K G W E

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I D I D S L E S I A D K N T T A M V I I

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N P N N P C G S V Y S Y D H L A K

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V A E V A R K

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L G I L V I A D E V Y G K L V L G S A P

TTATCCCGATGGGCGTCTTTGGGCACATTGCCCCGGTCTTGTCATTGGATCTCTGTCCA
F I P M G V F G H I A P V L S I G S L S

AGTCGTGGATAGTGCCTGGATGGCGACTTGGATGGGTGGCGGTGTACGACCCACAAAGA
K S W I V P G W R L G W V A V Y D P T K

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I L E K T K

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E A L P K I L E N T K A D F F K R I I

TGGTCTACTAAAGGAATCATCAGAGATATGTTATAGGGAAATAAAGGAAAACAAATATAT 8400
G L L K E S S E I C Y R E I K E N K Y I

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T C P H K P E G S M F V M

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 V K L N L H L L

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 E E I H D D I N F C C K L A K E E S V I

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 L C P

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 G S V

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 L G M E N W V R I T F A C V P S S L Q D

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 G L E R V K S F C Q R N K K K N S I N G

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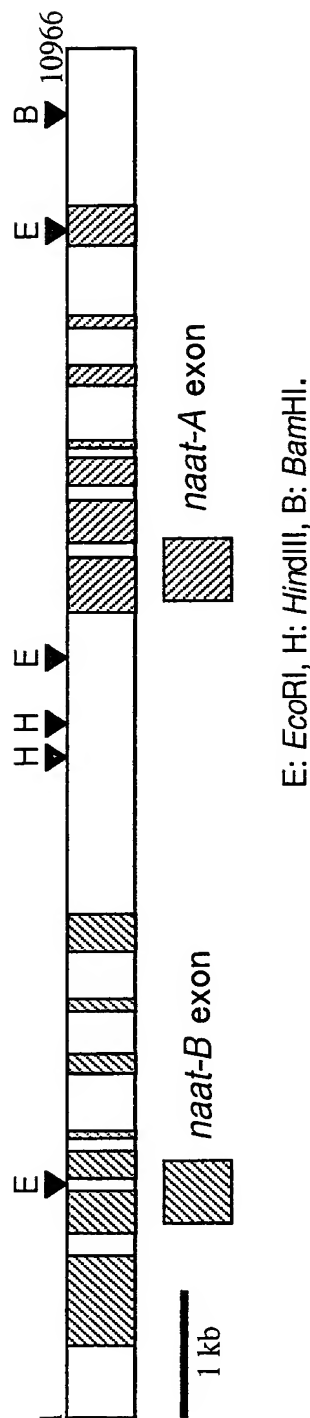
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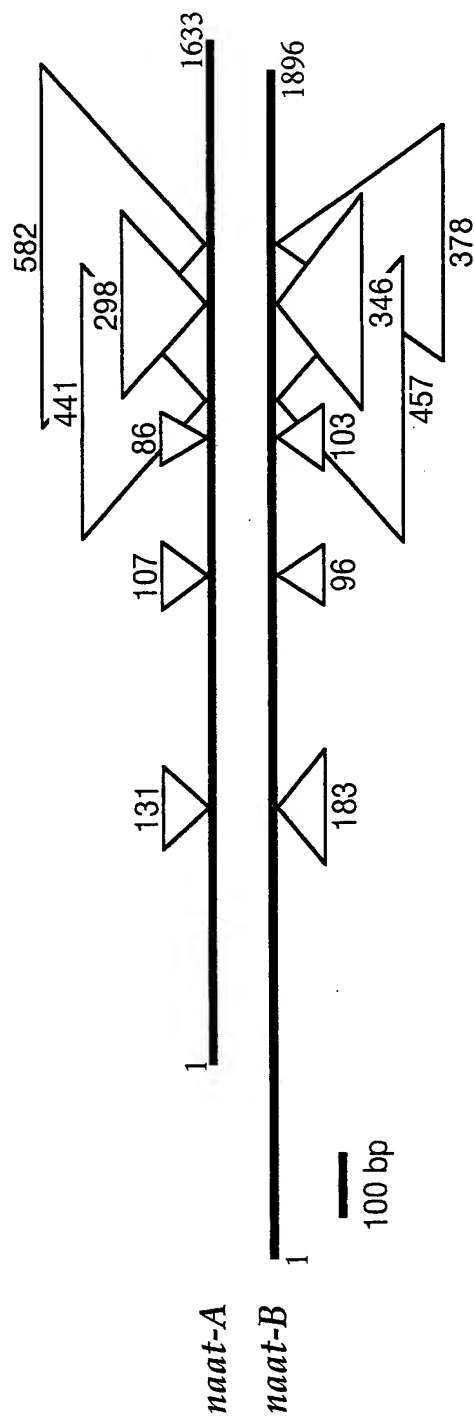
209240-EBZ6T00T

FIG. 11



A schematic view of the obtained genome fragment

FIG. 12



F I G. 1 3

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 IRAIRYKISASVEESGPRPVLPLAHGDPSVFPARTAVEAEDAVAAALRTGQFNCAAGV
 GLPAARSASVAEHL SQGV PYKLSADDVFLTAGGTQAIEV IIPVLAQTAGANILLPRPGYPN
 YEAAAAFNKLEVRHFDLIPDKGWEIDIDSLESIADKNTTAMVI INPNNPCGSVYSYDHLA
 KVAEVARKLGILVIADEVYGKLVLSAPFIPMGVFGHIAPVLSIGSLSKSWIVPGWRLGW
 VAVYDPTKILEKTKIST SITNYLNVSTDPATFVQEALPKILENTKADFFKRIIGLLKES
 EICYREIKENKYITCPHKPEGSMFVMVKLNLHLLLEEIHDDINFCKLAKEESVILCPGSV
 LGMENWVRITFACVPSSLQDGLERVKSFCQRNKKNSINGC*

F I G. 1 4

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 GKSNGHAEATANGHGEATANGKTNGHRESNGHAEADANGESNEHAEDSAANGESNGHAA
 AAAAAEEAVEWNFAGAKDGVLAATGANMSIRAIRYKISASVQEKGRPVLPALAHGDPSVF
 PAFRTAVEAEDAVAAALRTGQFNCAAGVGLPAARSASVAEHL SQGV PYMLSADDVFLTAG
 GTQAIEV IIPVLAQTAGANILLPRPGYPNYEAAAAFNRLVRFHFDLIPDKGWEIDIDSLE
 SIADKNTTAMVI INPNNPCGSVYSYDHL SKVAEVAKRLGILVIADEVYGKLVLSAPFIP
 MGVFGHITPVLSIGSLSKSWIVPGWRLGWVAVYDPRKILQETKIST SITNYLNVSTDPAT
 FIQAALPQILENTKEDFFKAIIGLLKESSEICYKQIKENKYITCPHKPEGSMFVMVKLNL
 HLLLEEIDDDIDFCCKLAKEESVILCPGSVLGMANWVRITFACVPSSLQDGLGRIKSFCQR
 NKKRNSSDDC*

FIG. 15 A



FIG. 15 B



FIG. 16

